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IN THE SPECIFICATION:

Please replace the paragraph beginning at page 1, line 3 with the following rewritten paragraph:

This application claims the benefit of U.S. Previsional Application No. 60/092,438, filed July 10, 1998, new abandoned and is a divisional application of U.S. Application No. 09/347,801 filed July 2, 1999, and issued on July 17, 2001 as now U.S. Patent 6,262,345, which claims the benefit of U.S. provisional Application No. 60/092,438, filed July 10, 1998.

Please replace the paragraph at page 2, lines 36-38 with the following rewritten paragraph:

Figures 1A. 1B. 1C and 1D shows a comparison of the amino acid sequences set forth in SEQ ID NOs:2, 4, 6 and 8 and the Zea mays and Arabidopsis thaliana sequences (SEQ ID NOs:17 (gi 3320104), 18 (gl 1839597) and 19 (gl 3402722)).

Please replace the paragraph at page 3, lines 1-3 with the following rewritten paragraph:

Figures 2A, 2B and 2C shows a comparison of the amino acid sequences set forth in SEQ ID NOs:10, 12, 14 and 16 the *Arabidopsis thaliana* and *Medicago sativ* sequences (SEQ ID NOs:21 (gl 1170711), 22 (gl 1709129), 23 (gl 1709127) and 24 (gl 1480078)).

Please replace the paragraph at page 19, lines 2-6 with the following rewritte paragraph:

Figures 1A-1D presents an alignment of the amino acid sequences set forth in SEQ ID NOs:2, 4, 6 and 8 and the Zee mays and Arabidopsis thaliana sequences (SEQ ID NOs:17 (gli 3320104), 18 (gli 1839597) and 19 (gli 3402722)). The data in Table 4 represents a calculation of the percent identity of the amino acid sequences set forth in SEQ ID NOs:2, 4, 6 and 8 and the Zee mays and Arabidopsis thaliana sequences (SEQ ID NOs:17, 18 and 19).

Please replace the paragraph at page 20, lines 5-10, underneath Table 5 with the following paragraph:

Figures 2A-2C presents an alignment of the amino acid sequences set forth SEQ ID NOs:10, 12, 14 and 16 the *Arabidopsis thaliana* and *Medicago sativa* sequences (SEQ ID NOs:20 (gi 1170711), 21 (gi 1709129), 22 (gi 1709127) and 23 (gi 1460078)). The data in Table 6 represents a calculation of the percent identity of

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the amino acid sequences set forth in SEQ ID NOs:10, 12, 14 and 16 and the Arabidopsis thaliana and Medicago sativa sequences (SEQ ID NOs: 20, 21, 22 and 23).

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